

Romany Population Data on Two Loci (D1S80, Apo BII) from East Slovakia

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ABSTRACT The present paper informs about allele frequency distributions for the two VNTR loci (D1S80, ApoBII) in East Slovak-Romanies. The results are compared with data of non-Romany Slovak population. Twelve segregating alleles were detected in a total sample of 87 Romany individuals. The most frequent alleles were 37 and 35 on the locus for ApoBII gene. 37/37 occurs as the most frequent genotype in East Slovak-Romanies. 13 alleles were detected in a total sample of 46 Romany individuals on the locus D1S80. 24 and 18 alleles and 24/24 genotype were the most frequent in Romany individuals.

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